

Epigenetic Re-Expression of Hemoglobin F Using Reversible LSD1 Inhibitors - Potential Therapies for Sickle Cell Disease

Steven Holshouser,¹ Rebecca Cafiero,¹ Mayra Robinson,¹ Joy Kirkpatrick,¹ Robert A. Casero,² Jr., Hyacinth I. Hyacinth³ and Patrick M. Woster.^{1*}

¹Department of Drug Discovery and Biomedical Sciences, Medical University of South Carolina, 70 President St., Charleston, SC 29414, USA; ²Sidney Kimmel Comprehensive Cancer Center, Johns Hopkins School of Medicine, 1650 Orleans St. Room 551, Baltimore, MD 21287, USA; ³Department of Pediatrics, School of Medicine, Emory University, 2015 Uppergate Dr., Atlanta, GA 30322, USA.

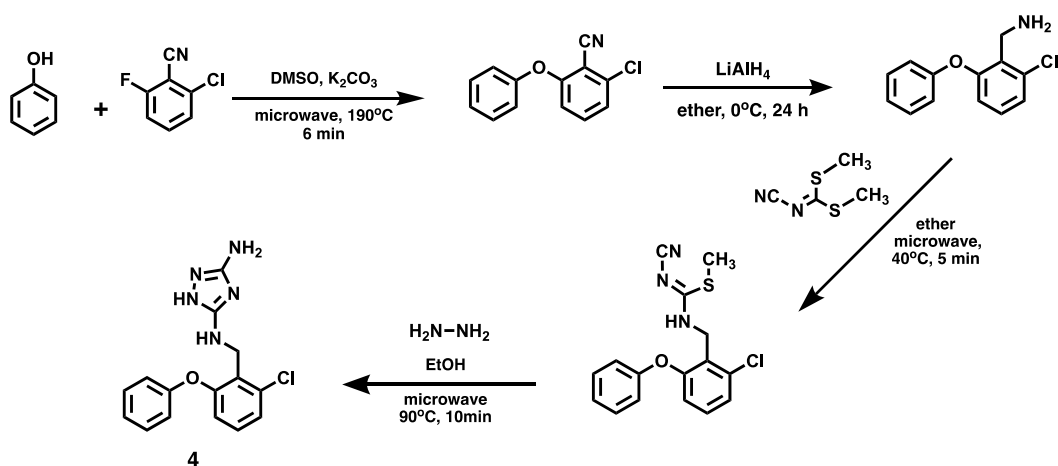
KEYWORDS: Epigenetics; lysine-specific demethylase 1; hemoglobin F; sickle cell disease.

Supporting Information

Synthesis (all compounds in this manuscript have been previously reported)

All reagents and dry solvents were purchased from Aldrich Chemical Co. (Milwaukee, WI), Sigma Chemical Co. (St. Louis, MO), VWR (Radnor, PA) or Fisher Scientific (Chicago, IL) and were used without further purification except as noted below. Triethylamine was distilled from potassium hydroxide and stored in a nitrogen atmosphere. Dry methanol, ethyl acetate, tetrahydrofuran, dimethyl formamide and hexane were prepared using a Glass Contour Solvent Purification System (Pure Process Technology, LLC, Nashua, NH). Preparative scale chromatographic procedures were carried out using a CombiFlash Rf200 chromatography system (Teledyne-Isco, Lincoln, NE) fitted with silica gel 60 cartridges (230-440 mesh). Thin layer chromatography was conducted on Merck precoated silica gel 60 F-254. Ion exchange chromatography was conducted on Dowex1X8-200 anion exchange resin.

All ¹H- and ¹³C-NMR spectra were recorded on a Bruker Avance 600 MHz spectrometer, and all chemical shifts are reported as δ values referenced to TMS or DSS. Splitting patterns are indicated as follows: s, singlet; d, doublet; t, triplet; m, multiplet; br, broad peak. In all cases, ¹H-NMR, ¹³C-NMR and MS spectra were consistent with assigned structures. Mass spectra were recorded by LC/MS on a Waters autopurification liquid chromatography with a model 3100 mass spectrometer detector. Prior to biological testing procedures, all compounds were determined to be >95% pure by UPLC chromatography (95% H₂O/5% acetonitrile to 20% H₂O/80% acetonitrile over 10 minutes) using a Waters Acquity H-series ultrahigh-performance liquid chromatograph fitted with a C18 reversed-phase column (Acquity UPLC BEH C18 1.7 M, 2.1 X 50 mm). Compounds **4-11** were synthesized according to the general procedures described below.



Scheme S1. Synthetic route to 3,5-diamino-1,2,4-triazole analogues **4-6**.

Method A¹

General procedure for the synthesis of substituted 2-chloro-6-aryloxybenzonitriles

Synthesis of 2-chloro-6-phenyloxybenzonitrile. To a 20 mL microwave vial containing a magnetic stir bar was added 0.78 g (5.0 mmol) of 2-fluoro-6-chlorobenzonitrile, (1.04 g, 7.5 mmol) of K₂CO₃, 0.52 g (5.5 mmol) of phenol and 12.0 mL of anhydrous DMSO. The vial was then sealed and stirred to distribute the contents evenly. The mixture was then microwaved at 190°C for 6 min at high absorption to insure even heating. The reaction mixture was poured into a beaker containing 100 mL of crushed ice to precipitate the product. The aqueous layer was extracted with three 50 mL portions of diethyl ether, and the ether layer was washed with 25 mL of saturated NaCl, dried over anhydrous Na₂SO₄, filtered, and the ether was removed in vacuo to yield 1.11 g of the desired diaryl ether (97% yield). The crude product was pure enough to be used in the next reaction without further purification.

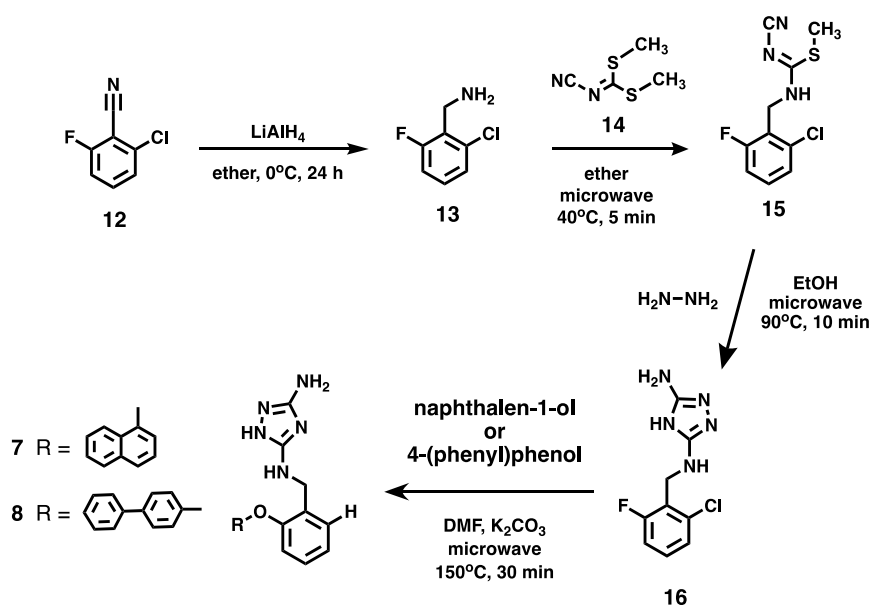
General procedure for the synthesis of substituted 2-chloro-6-aryloxybenzylamines

Synthesis of 2-chloro-6-phenyloxybenzylamine. A 1.11 g portion of 2-chloro-6-phenoxybenzonitrile (4.8 mmol) was dissolved in 50 mL of anhydrous diethyl ether, cooled to 0°C in an ice bath and stirred while bubbling dry argon into the reaction mixture for 10 min. A 14.49 mL portion of 1.0 M LiAlH₄ in THF (14.49 mmol) was then added dropwise with stirring over 20 min. The resulting reaction mixture was allowed to stir for 2 hrs at 0°C, and then warmed to room temperature and allowed to stir overnight. The mixture was cooled to 0°C, and the reaction was quenched by the slow addition of Na₂SO₄•10 H₂O. When the evolution of gas subsided, the reaction was stirred for 30 min at room temperature, and the mixture was filtered through a Celite pad. The filtrate was concentrated to dryness to yield crude 2-chloro-6-benzoyloxybenzylamine. The crude product was pure enough to be used in the next reaction without further purification.

General procedure for the synthesis of substituted N³-(2-chloro-6-aryloxybenzyl)-1H-1,2,4-triazole-3,5-diamines

Synthesis of N³-(2-chloro-6-phenoxybenzyl)-1H-1,2,4-triazole-3,5-diamine, 4. A 0.935 g portion of 2-chloro-6-phenoxybenzyl amine (4.0 mmol) was dissolved in 12 mL of diethyl ether and added to a 20 mL microwave vial equipped with a magnetic stir bar. A 0.702 g portion of dimethyl cyanodithioiminocarbonate (4.8 mmol) was added and the vial was sealed. The contents were microwaved at 45°C for 5 min, cooled to room temperature, and the ether was removed in vacuo to yield the intermediate as a white to pale yellow solid. A 0.192 g portion of hydrazine hydrate (6.0 mmol) in 12 mL of dry ethanol was then injected, the vial was stirred to break up the solid intermediate, and the resulting mixture was microwaved at 90°C for 10 min at high absorption. The ethanol was removed in vacuo, and the residue was purified by silica gel chromatography (9% MeOH in CH₂Cl₂) to afford 1.07 g of pure **4** (85%) as an off-white, amorphous solid. ¹H-NMR (400MHz, CD₃OD/TMS) δ 4.21 (s, 2H), 6.77-6.80 (dd, 1H), 6.99-7.01 (d, 2H), 7.11-7.15 (t, 1H), 7.20-7.27 (m, 2H), 7.33-7.38 (t, 2H). UPLC retention time: 12.1 min. MS calculated 315.09, found 316.33 ([M+1]⁺)

Method B²



Scheme S2. Synthetic route to dechlorinated compounds **7** and **8**.

Synthesis of 2-chloro-6-fluorobenzylamine, 13. 2-chloro-6-fluorobenzonitrile **12** (1.0 g, 6.43 mmol) was dissolved in 50 mL of diethyl ether and stirred at 0°C while bubbling argon through the solution for 10 minutes, followed by the dropwise addition of 15 mL (3.0 equivalents) of 1.0 M LiAlH₄ in THF over 20 min. Following addition of the reducing agent, the reaction was stirred for 2 hours at 0°C, warmed to room temperature and allowed to stir overnight. The reaction was cooled to 0°C and quenched by slow addition of Na₂SO₄•10 H₂O. When the evolution of gas was complete, the reaction was warmed to room temperature and allowed to stir for an additional 30 minutes, and then filtered through a Celite pad. The filtrate was then concentrated to dryness *in vacuo*. The crude material was purified on silica (0-5% MeOH in dichloromethane), to produce the desired 2-chloro-6-fluorobenzylamine **13** (0.913 g, 89% yield). ¹H NMR (400 MHz, CDCl₃) δ 4.88 (s, 2H), 7.09 (m, 1H), 7.28 (m, 2H).

Synthesis of N³-(2-chloro-6-fluorobenzyl)-1H-1,2,4-triazole-3,5-diamine, 16. A 0.913 g (5.72 mmol) of 2-chloro-6-fluorobenzylamine **13** was dissolved in 12 mL of diethyl ether and added to a 20 mL microwave vial equipped with a magnetic stirring bar. Dimethyl cyanodithioiminocarbonate **14** (1.0 g, 6.43 mmol) was then added and the vial was sealed. The contents were microwaved at 45°C for 5 minutes, then allowed to cool to room temperature, and the ether was removed *in vacuo* to yield the intermediate **15** as a white solid. Hydrazine hydrate (0.275 g, 8.58 mmol) in 12 mL of ethanol was then injected into the vial, and the reaction was stirred to break up the intermediate solid. The dispersed reaction mixture was microwaved at 90°C for 10 min on high absorption. The mixture was allowed to cool to room temperature, and the ethanol was removed *in vacuo* to yield the crude product. The crude material was purified on silica (10% MeOH in dichloromethane), to afford 1.27 g of pure N³-(2-chloro-6-fluorobenzyl)-1H-1,2,4-triazole-3,5-diamine **16** as a white, amorphous powder (92% yield). ¹H NMR (400 MHz, CDCl₃) δ 4.32 (s, 2H), 7.20-7.22 (t, 4H), 7.31-7.36 (m, 2H). UPLC retention time: 0.46 min. MS calculated 241.05, found 242.30 ([M+1]⁺)

Synthesis of N³-(2-(naphthalen-1-yloxy)benzyl)-1H-1,2,4-triazole-3,5-diamine, 7. To a 20 mL microwave vial equipped with a magnetic stir bar was added N³-(2-chloro-6-fluorobenzyl)-1H-1,2,4-triazole-3,5-diamine **16** (1.0 g, 4.14 mmol) (0.86 g, 6.21 mmol) K₂CO₃, naphthalen-1-ol (0.659 g, 4.55 mmol), and 12 mL of anhydrous dimethyl sulfoxide. The vial was then sealed and stirred to distribute contents evenly. The reaction was microwaved at 190°C for 6 min on high absorption, and the contents were poured into a beaker containing 100 mL crushed ice. The product was extracted with three 50 mL portions of diethyl ether, and the ether layer was washed with 25 mL of saturated NaCl. The organic layer was dried over Na₂SO₄, filtered, and the filtrate was removed *in vacuo* to provide crude **7**. The crude material was purified on silica (10% MeOH in dichloromethane), to afford pure **7** in 56% yield as a white solid. ¹H NMR (400 MHz, CDCl₃) δ 4.66 (s, 2H), 7.36-7.38 (d, 2H), 7.53-7.57 (d, 1H), 7.63-7.64 (m, 1H), 7.68-7.70 (d, 1H), 7.84-7.85 (d, 1H), 7.91-7.93 (d, 2H), 8.15-8.16 (m, 2H), 8.17-8.19 (m, 1H). UPLC retention time: 0.44 min. MS calculated 331.14, found 332.43 ([M+1]⁺)

Synthesis of N³-[6-(4-phenyl)phenyloxy]-1H-1,2,4-triazole-3,5-diamine 8. To a 20 mL microwave vial equipped with a magnetic stir bar was added N³-(2-chloro-6-fluorobenzyl)-1H-1,2,4-triazole-3,5-diamine (1.0 g, 4.14 mmol) (0.86 g, 6.21 mmol) K₂CO₃, 4-phenylphenol (0.715 g, 4.55 mmol), and 12 mL of anhydrous dimethyl sulfoxide. The vial was then sealed and stirred to distribute contents evenly. The reaction was microwaved at 150°C for 6 min on high absorption, and the contents were poured into a beaker containing 100 mL crushed ice. The product was extracted with three 50 mL portions of diethyl ether, and the ether layer was washed with 25 mL of saturated NaCl. The organic layer was dried over Na₂SO₄, filtered, and the filtrate was removed *in vacuo* to provide crude **8**. The crude material was purified on silica (10% MeOH in dichloromethane), to afford 1.14 g of pure **8** in 64% yield. ¹H NMR (400 MHz, CD₃OD) δ 4.74 (s, 2H), 6.85-6.87 (d, 1H), 7.26-7.28 (d, 2H), 7.38-7.42 (m, 1H), 7.47-7.51 (m, 2H), 7.62-7.64 (d, 2H), 7.68-7.71 (d, 2H), 7.77-7.81 (m, 1H), 7.85-7.88 (m, 1H), 8.15-8.17 (d, 1H). UPLC retention time: 0.43 min. MS calculated 357.16, found 358.37 ([M+1]⁺) Note: This reaction yielded a dechlorinated product due to stabilization of the aromatic ring via anchimeric assistance of the adjacent secondary nitrogen. The stabilization of the aromatic ring increases the ability of chlorine to act as a good leaving group, resulting in hydrogen abstraction and formation of the dechlorinated product.

Peptide Synthesis^{3,4}

N^α-Fmoc amino acids were purchased from the Advanced Chemtech (Louisville, KY) and AAPPTec (Louisville, KY). Fmoc-rink linker and aminomethylated polystyrene resin was purchased from the Novabiochem (Gibbstown, NJ). Reagent grade Piperidine was purchased from the Sigma Aldrich. All the other solvents were purchased from the VWR and Fisher and used without further purification. All of the cyclic and linear peptide analogues were synthesized by using standard N^α-Fmoc/tBu solid-phase peptide synthesis. Three channel PS3 automated peptide synthesizer from Protein Technologies, Inc. Tucson Arizona was used for peptide synthesis. The aminomethylated polystyrene resin (0.25 mmol, 0.36 mmol/g) was placed in a 40 mL glass reaction vessel in the synthesizer and allowed to swell in 15 mL of DMF solution for 30 min. Then the resin was washed with 15 mL of DMF (3 X 2min). The Fmoc linker was introduced to the swelled resin using mixture of

Fmoc linker (1.0 mmol, 4 equiv), HBTU (1 mmol, 4 equiv) and NMM (2 mmol, 8 equiv) in DMF for 60 min. After coupling of the Fmoc linker to the resin, the resin was washed with 15 mL of DMF solution (5 X 2 min). The Fmoc protecting group on the resin was removed with 15 mL of 20% piperidine in DMF (2 x 15 min) followed by washing with 15 mL of DMF (5 X 2 min). Then, a preactivated Fmoc-amino acid prepared by mixing a Fmoc-amino acid (4 equiv), HBTU (4 equiv), and NMM (8 equiv) in DMF was introduced into the reaction vessel, and the reaction was continued for 1 hour. The deprotection and coupling steps were repeated for each amino acid until desired sequence was obtained.

Once the fully protected peptide having desired sequence is obtained on resin (0.25 mmol), orthogonal protective groups of the peptide (alloc protecting groups of Lys and allyl group of Glu) were selectively removed using mixture of Pd(PPh₃)₄ (30 mg, 0.1 equiv), and DMBA (390 mg, 10 equiv) in 6 mL DMF:DCM (1:3) in the 40 mL reaction vessel under N₂ atmosphere for 30 min for two times. Then the resin was washed with 15 mL of DMF (5 X 2 min) and washed with 15 mL of 0.1 M LiCl in DMF solution (3 x 2 min). It was again washed with 15 mL of DMF (3 X 2 min). The resin was treated with PyBOP/HOBt/DIPEA (6, 6, and 12 equiv) in 6 mL of DCM: DMF: NMP (1:1:1) for 6 h twice for formation of the lactam bridge. Then resin was washed with DMF (3 X 2 min) and the cyclic peptide was cleaved from the resin as very similar to the procedure described above for the linear peptide.

The purified peptides were characterized by HRMS and LC-MS. High resolution mass spectrometric data was taken in the positive ion mode using Bruker AUTOFLEX III MALDI-TOF instrument. LC-MS data was obtained in the positive ion mode using Waters LC-MS instrument [having Waters 2545 quaternary gradient module, Waters 2767 sample manager, Waters SFO fluidic organizer, Waters 3100 mass detector containing single quadrupole, and Waters PDA detector 2998] on Waters Xterra C18 column (3.0 x 100 mm, 5 μM). UPLC chromatograms obtained using a Waters Acquity UPLC (H class, PDA detector, sample manager FTN and quaternary solvent manager) fitted with a Waters BEH C18 column (2.1 x 100 mm, 1.7 μM).

Analytical Data for Compounds 4-11 (from references 52-54)

*N*³-(2-chloro-6-phenoxybenzyl)-4*H*-1,2,4-triazole-3,5-diamine **4**. ¹H-NMR (400MHz, CD₃OD/TMS) δ 4.21 (s, 2H), 6.77-6.80 (dd, 1H), 6.99-7.01 (d, 2H), 7.11-7.15 (t, 1H), 7.20-7.27 (m, 2H), 7.33-7.38 (t, 2H). UPLC retention time: 12.1 min. MS calculated 315.09, found 316.33 ([M+1]⁺)

*N*³-(2-chloro-6-(naphthalen-1-yloxy)benzyl)-1*H*-1,2,4-triazole-3,5-diamine, **5**. ¹H NMR (400 MHz, DMSO) δ 4.49 (s, 2H), 6.67-6.80 (dd, 1H), 6.99-7.01 (d, 1H), 7.25 (d, 2H), 7.54-7.59 (m, 3H), 7.77-7.79 (dd, 1H), 7.99-8.01 (dd, 1H), 8.12-8.14 (dd, 1H). UPLC retention time: 14.34 min. MS calculated 365.10, found 366.21([M+1]⁺)

*N*³-(2-chloro-6-(naphthalen-2-yloxy)benzyl)-1*H*-1,2,4-triazole-3,5-diamine, **6**. ¹H NMR (400 MHz, CD₃OD) δ 2.14 (s, 2H), 6.72 (dd, 1H), 6.81-6.86 (dd, 1H), 7.07 (d, 1H), 7.08 (s, 1H), 7.17 (d, 1H), 7.32 (m, 3H), 7.51 (dd, 1H), 8.19 (dd, 1H). UPLC retention time: 14.53 min. MS calculated 365.82, found 366.21 ([M+1]⁺)

*N*³-(2-(naphthalen-1-yloxy)benzyl)-1*H*-1,2,4-triazole-3,5-diamine, **7**. ¹H NMR (400 MHz, CDCl₃) δ 4.66 (s, 2H), 7.36–7.38 (d, 2H), 7.53–7.57 (d, 1H), 7.63–7.64 (m, 1H), 7.68–7.70 (d, 1H), 7.84–7.85 (d, 1H), 7.91–7.93 (d, 2H), 8.15–8.16 (m, 2H), 8.17–8.19 (m, 1H). UPLC retention time: 0.44 min. MS calculated 331.14, found 332.43 ([M + 1]⁺).

*N*⁵-(2-([1,1'-biphenyl]-4-yloxy)benzyl)-1*H*-1,2,4-triazole-3,5-diamine, **8**. ¹H NMR (400 MHz, CD₃OD) δ 4.74 (s, 2H), 6.85–6.87 (d, 1H), 7.26–7.28 (d, 2H), 7.38–7.42 (m, 1H), 7.47–7.51 (m, 2H), 7.62–7.64 (d, 2H), 7.68–7.71 (d, 2H), 7.77–7.81 (m, 1H), 7.85–7.88 (m, 1H), 8.15–8.17 (d, 1H). UPLC retention time: 0.43 min. MS calculated 357.16, found 358.37 ([M + 1]⁺).

H-ARAM-*c*[K⁵TARKE¹⁰]TGG-KAPRKQLAK(*N*-CH₃-(CH₂)₆CO)-NH₂, **9**. UPLC retention time: 0.273 min; MS for C₁₁₁H₂₀₃N₃₇O₂₉S: Calc. 2550.53, found 845.6 ([M+3H])⁺.

H-ARTM-*c*[K⁵TARKE¹⁰]TGG-KAPRKQLAK(*N*-CH₃-(CH₂)₆CO)-NH₂, **10**. UPLC retention time: 0.269 min; MS for C₁₁₂H₂₀₅N₃₇O₃₀S: Calc. 2580.54, found 855.5 ([M+3H])⁺.

H-ARAM-*c*[K⁵TARKE¹⁰]TGG-KAPRKQLA-OH, **11**. UPLC retention time: 5.16min; HRMS for C₉₅H₁₇₃N₃₇O₂₄S: Calc. 2250.67, found 2251.357 (M+1H)⁺

Enzyme Assay

Compounds were evaluated for the ability to inhibit recombinant LSD1/CoREST using a commercially available assay kit (#700120, Cayman Chemical, Ann Arbor, MI). The substrate and all compounds were incubated in assay buffer from 30 min up to 4 h at 37 °C as described in the commercial protocol. The volume of each reaction well

was 50 μ L, containing 5 μ L of a 200 μ M solution of substrate peptide and 20 μ L of a 15 ng/ μ L enzyme solution. All compounds were diluted in 1% DMSO with assay buffer to a final volume of 50 μ M. Fluorescence was measured at the recommended wavelengths of k_{ex} = 530 nm, k_{em} = 590 nm. IC₅₀ determinations were performed using serial dilutions at 250, 125, 62.5, 31.25, 6.25, 3.125, 1.563, 0.781 and 0.390 μ M. All blanks contained 1% DMSO to determine any solvent effects.

Cell Culture

K562 (human myelogenous leukemia) cells were purchased from ATCC. Cells were cultured in Iscove's Modified Dulbecco's Medium containing 10% (v/v) fetal bovine serum and 5% penicillin and streptomycin, as previously described.⁵⁴ All cultures were grown at 37°C in a humidified environment containing 5% CO₂.

CD34⁺ cells were expanded in H3000 media supplemented with CC100 (Stem Cell Technologies) for 4 days at a density of 10⁵ cells/mL prior to use for compound evaluation. Cells were treated for 24, 48 or 72 hours with an appropriate concentration of each analogue.

Cell Viability Assay

All cell lines were purchased from ATCC (Manassas, VA). For the (3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxy-methoxy-phenyl)-2-(4-sulfo-phenyl)-2H-tetrazolium) (MTS) reduction assay, 2000 cells per well were seeded in 50 μ L of complete medium per well of a 96-well plate and the cells were allowed to attach overnight at 37°C in a 5% CO₂ atmosphere. The medium was aspirated and cells were treated with 100 μ L of fresh medium containing appropriate concentrations of each inhibitor to be tested. The cells were incubated for 72 h at 37°C in 5% CO₂, after which 20 μ L of the MTS reagent solution (Promega CellTiter 96 Aqueous One Solution Cell Proliferation Assay) was added to the medium. The cells were incubated for another 2 h at 37°C, and absorbance was measured at 490 nm on a SpectraMax M5 instrument (Molecular Devices) equipped with SOFTmax PRO 4.0 software to determine cell viability. A reference wavelength of 690 nm was used to subtract the background. Percent cell death was calculated by the following equation: % Cell Death = (Abs Control - Abs sample)/Abs Control X100. A dose response curve was constructed for each inhibitor, and each data point was the average of 3 determinations obtained during a single experiment \pm S.E.M. IC₅₀ values were calculated using the GraphPad Prism 5 software package (Graph-Pad, San Diego, California).

RT-qPCR

Media was rinsed with PBS and cells were lysed using TriZol Reagent (Invitrogen, Cat# 15596026). mRNA was isolated according to manufacturer's protocols and purity confirmed using a Nanodrop-1000 spectrophotometer (Thermo Fisher). Quantitative reverse-transcription real time polymerase chain (qRT-PCR) reaction was run using a High-Capacity cDNA Reverse Transcription Kit (Applied Biosystems, Cat# 4368814) followed by TaqMan® Fast Advanced Master Mix (Applied Biosystems, Cat# 4444557) with TaqMan® Gene Expression Assay Primers (Applied biosystems, listed below) using a StepOne Plus instrument (Thermo Fisher). TNF- α , IL-6 and the internal control GAPDH were then quantitated for each sample in triplicate. Results are reported as fold change ($2^{-\Delta\Delta CT}$).

Primary rat cardiomyocyte isolation

Primary male Sprague-Dawley rat cardiomyocytes were isolated via a hanging heart preparation using enzymatic digestion, as previously described.⁵² In brief, rats were euthanized with 5% isoflurane vaporized in 100% O₂. The heart was retrogradely perfused with collagenase. The cardiomyocytes were plated on 6-well culture trays that were coated with laminin at an initial plating density of 1.5×10^5 cells/well. After overnight incubation, the cardiomyocytes were rinsed and maintained in serum-free medium.

Co-immunoprecipitation

Primary rat cardiomyocytes were isolated as described earlier and treated with varying concentrations of test compounds. Cells were lysed and scraped with IP Lysis Buffer (20mM Tris-Cl, 150mM NaCl, 1mM EDTA, 1mM EGTA, 1mM β -glycerol, 2.5mM Na pyrophosphate). Dynabeads (Life Biosciences) were added (1:10 beads:lysate v/v) and incubated at 4°C for 1-hour to pre-clear. Mixture was centrifuged (5000rpm) and the supernatant was saved. An antibody (2-5 μ g) for the protein of interest was added and rocked overnight in cold room. Dynabeads (20 μ L) were added and incubated at room temp for 1-hour. Centrifuge mixture for 1-minute at 5000rpm. The supernatant was decanted and the pellet washed 3X with IP Lysis Buffer with 0.1% Triton X-100. Samples were then analyzed by SDS-PAGE and immunoblotted.

Preliminary Toxicology

Toxicology screen of Compound 4 in CD1 mice: We performed a pilot toxicology study in four CD1 mice to determine whether 3,5-diamino-1,2,4-triazoles such as **4** caused any acute organ toxicities as determined by blood factor analysis and histology. This study was performed in collaboration with the MUSC Veterinary Services. In short, a 3mg/kg, intraperitoneal injection of compound **4** was administered daily for 1-week. After the drug course, mice were evaluated by the MUSC veterinarian. In vivo treatment with **4** revealed only minor anomalies: slight elevation of the BUN/creatinine ratio and minor thrombocytopenia. Histology revealed minor ectatic cortical tubules with attenuated epithelium and minor vacuolization of cardiomyocytes. No hepatic inflammation or necrosis, lung pathology or splenic changes were seen. A complete blood panel, liver enzymes, and kidney functional labs were obtained, followed by histological evaluation of various organs (i.e. heart, lung, liver, spleen, and kidney). A comprehensive report on histological damage was obtained by a veterinary pathologist. In addition, a consultation with an MUSC Hematology Resident and Nephrologist were also performed. Upon gross examination, all four mice showed no abnormalities or contraindications from drug treatments. The blood panel and liver enzymes had no outstanding findings and were within normal range. However, blood urea nitrogen (BUN) was elevated, creatinine was low, and blood glucose was elevated (**Table S1**). Thus, the BUN:creatinine ratio was higher than reported normal values. In addition, the CBC showed minor thrombocytopenia.

Table S1. Laboratory values for CD-1 mice treated for 1 week with compound **4**.

Blood Diagnostic Tests			
	Normal Range #	Compound 9 ‡	
ALT	45.08 ± 2 u/L	42.75 ± 5.0	
AST	80.55 ± 8.3 u/L	165.25 ± 17.8	
ALP	14-118 u/L	66.67 ± 21.5	
CREATININE	11.3 ± 0.1 mg/dL	0.15 ± 0.03	<i>L</i>
BUN	14.68 ± 0.5 mg/dL	22 ± 1.5	<i>H</i>
GLU	164.89 ± 3.5 mg/dL	263.5 ± 19.5	<i>H</i>
<i>Leukocytes</i>			
WBC	8.79 ± 0.3 (k/μL)	5.3 ± 0.9	
NE	1.5 ± 0.1 (k/μL)	1.1 ± 0.1	
LY	6.59 ± 0.3 (k/μL)	4.0 ± 0.8	
MO	0.49 ± 0.03 (k/μL)	0.13 ± 0.03	<i>L</i>
EO	0.17 ± 0.02 (k/μL)	0.07 ± 0.02	
BA	0.04 ± 0.01 (k/μL)	0.01 ± 0.01	
NE%	6.6-38.9%	20.5 ± 2.1	
LY%	55.8-91.6%	75.3 ± 2.6	
MO%	0-7.5%	2.5 ± 0.6	
EO%	0-3.9%	1.4 ± 0.5	
BA%	0-2%	0.31 ± 0.2	
<i>Erythrocytes</i>			
RBC	8.93 ± 0.2 (M/μL)	10.3 ± 0.9	
Hb	14.67 ± 0.3 (g/dL)	15.6 ± 1.4	
HCT	49.97 ± 1.1 (%)	54.4 ± 5.3	
MCV	56.1 ± 0.6 (fL)	52.8 ± 1.8	
MCH	16.45 ± 0.6 (pg)	15.2 ± 0.3	
MCHC	29.5 ± 0.4 (g/dL)	28.9 ± 1.0	
RDW	16.79 ± 0.1 (%)	16.4 ± 0.4	
<i>Thrombocytes</i>			
PLT	1529 ± 53 (K/μL)	836.5 ± 324	<i>L</i>
MPV	5.1 ± 0.06 (%)	5.1 ± 0.1	

Normal clinical laboratory values as reported by Charles River Laboratories, CD1 mouse supplier

‡ 3mg/kg, i.p. QDx7days; n=4 CD1 mice

Histological evaluation by hemotoxylin and eosin staining of various organs revealed very minimal to minor pathological changes. Of note, the kidney showed minor ectatic cortical tubules with attenuated epithelium (**Figure S1A**). In addition, the heart showed minor vacuolation of cardiomyocytes without any interstitial immune infiltration (**Figure S1B**). There was also negligible multifocal hepatic inflammation and necrosis (**Figure S1C**). Finally, the spleen showed no outstanding lesions (not shown).

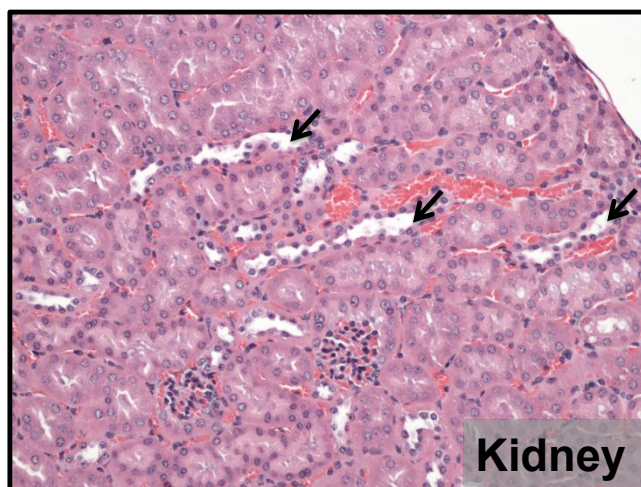


Figure S1A. Kidney histology of compound 4 treated CD1 male mice (3mg/kg, *i.p.*, QD 7 days). Hemotoxylin and eosin staining of cortical kidney. Mild multifocal ectatic tubules with attenuated epithelium (Arrow). Limited tubular hyperplasia and regeneration. (Reported in 2 of 4 animals).

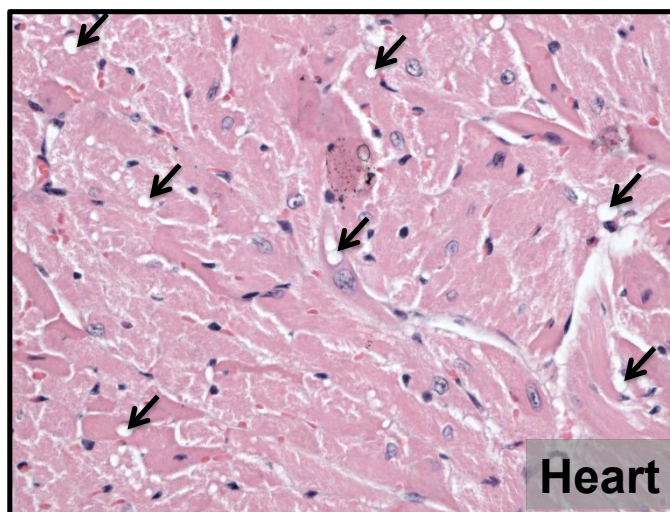


Figure S1B. Heart histology of compound 4 treated male CD1 mice (3mg/kg, *i.p.*, QD 7 days). Hemotoxylin and eosin staining of left ventricular free wall. Mild diffuse cardiomyocyte intrasarcoplasmic vacuolation without interstitial infiltrate. (Reported in 2 of 4 animals).

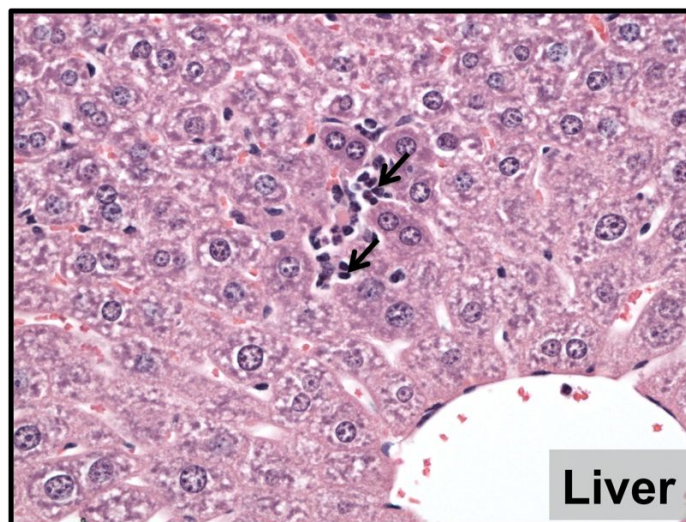


Figure S1C. Liver histology of compound 4 treated male CD1 mice (3mg/kg, *i.p.*, QD 7 days). Hemotoxylin and eosin staining of left lobe. Mild multifocal hepatic inflammation and necrosis can be found, as indicated by the arrows. (Reported in 3 of 4 animals).

References

52. Kutz, C. J.; Holshouser, S. L.; Marrow, E. A.; Woster, P. M.: 3,5-Diamino-1,2,4-triazoles as a novel scaffold for potent, reversible LSD1 (KDM1A) inhibitors. *MedChemComm* **2014**, *5* (12), 1863-1870. PM CID: 4286191.
53. Holshouser, S.; Dunworth, M.; Murray-Stewart, T.; Peterson, Y. K.; Burger, P.; Kirkpatrick, J.; Chen, H. H.; Casero, R. A., Jr.; Woster, P. M.: Dual inhibitors of LSD1 and spermine oxidase. *MedChemComm* **2019**, *10* (5), 778-790. PM CID: 6533887.
54. Kumarasinghe, I. R.; Woster, P. M.: Cyclic peptide inhibitors of lysine-specific demethylase 1 with improved potency identified by alanine scanning mutagenesis. *Eur. J. Med. Chem.* **2018**, *148*, 210-220. PM CID: PMC5837957.